

AMENDMENTS TO THE CLAIMS

1-3. (Canceled).

4. (Currently Amended) ~~The An isolated polypeptide of Claim 1~~ having at least 95% amino acid sequence identity to:

(a) the amino acid sequence of the polypeptide of shown in Figure 136 (SEQ ID NO: 136);

(b) the amino acid sequence of the polypeptide of shown in Figure 136 (SEQ ID NO: 136), lacking its associated signal peptide; or

~~(c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 136 (SEQ ID NO: 136);~~

~~(d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 136 (SEQ ID NO: 136), lacking its associated signal peptide; or~~

(e) (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203547;

wherein said isolated polypeptide is more highly expressed in normal esophagus tissue compared to esophageal tumor, or wherein said isolated polypeptide is encoded by a polynucleotide that is more highly expressed in normal esophagus tissue compared to esophageal tumor.

5. (Currently Amended) The isolated polypeptide of ~~Claim 1~~ Claim 4 having at least 99% amino acid sequence identity to:

(a) the amino acid sequence of the polypeptide of shown in Figure 136 (SEQ ID NO: 136);

(b) the amino acid sequence of the polypeptide of shown in Figure 136 (SEQ ID NO: 136), lacking its associated signal peptide;

~~(c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 136 (SEQ ID NO: 136);~~

~~(d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 136 (SEQ ID NO: 136), lacking its associated signal peptide; or~~

(e) (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203547;

wherein said isolated polypeptide is more highly expressed in normal esophagus tissue compared to esophageal tumor, or wherein said isolated polypeptide is encoded by a polynucleotide that is more highly expressed in normal esophagus tissue compared to esophageal tumor.

6. (Currently Amended) An isolated polypeptide comprising:
 - (a) the amino acid sequence of the polypeptide of shown in Figure 136 (SEQ ID NO: 136);
 - (b) the amino acid sequence of the polypeptide of shown in Figure 136 (SEQ ID NO: 136), lacking its associated signal peptide;
 - (c) ~~the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 136 (SEQ ID NO: 136);~~
 - (d) ~~the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 136 (SEQ ID NO: 136), lacking its associated signal peptide; or~~
 - (e)(c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203547.
7. (Currently Amended) The isolated polypeptide of Claim 6 comprising the amino acid sequence of the polypeptide of shown in Figure 136 (SEQ ID NO: 136).
8. (Currently Amended) The isolated polypeptide of Claim 6 comprising the amino acid sequence of the polypeptide of shown in Figure 136 (SEQ ID NO: 136), lacking its associated signal peptide.
9. (Canceled)
10. (Canceled)
11. (Original) The isolated polypeptide of Claim 6 comprising the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203547.
12. (Currently Amended) A chimeric polypeptide comprising a polypeptide according to ~~Claim 1~~ Claim 4 fused to a heterologous polypeptide.
13. (Currently Amended) The chimeric polypeptide of Claim 12, wherein said heterologous polypeptide is ~~an epitope~~ a tag polypeptide or an Fc region of an immunoglobulin.

14. (New) An isolated polypeptide having at least 95% amino acid sequence identity to:

(a) the amino acid sequence of the polypeptide of SEQ ID NO: 136;

(b) the amino acid sequence of the polypeptide of SEQ ID NO: 136, lacking its associated signal peptide;

(c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203547;

and wherein said isolated polypeptide or a fragment thereof can be used to generate an antibody which can be used to specifically detect the polypeptide of SEQ ID NO: 136 in esophagus tissue samples.

15. (New) The isolated polypeptide of Claim 14 having at least 99% amino acid sequence identity to:

(a) the amino acid sequence of the polypeptide of SEQ ID NO: 136;

(b) the amino acid sequence of the polypeptide of SEQ ID NO: 136, lacking its associated signal peptide;

(c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203547;

and wherein said isolated polypeptide or a fragment thereof can be used to generate an antibody which can be used to specifically detect the polypeptide of SEQ ID NO: 136 in esophagus tissue samples.

16. (New) A chimeric polypeptide comprising a polypeptide according to Claim 14 fused to a heterologous polypeptide.

17. (New) The chimeric polypeptide of Claim 16, wherein said heterologous polypeptide is a tag polypeptide or an Fc region of an immunoglobulin.

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DELETION OF INVENTORS

Please correct the inventorship under 37 C.F.R. § 1.48(b) by removing the following inventors from the present application:

Dan L. Eaton, Ellen Filvaroff, Mary E. Gerritsen, and Colin K. Wantanabe.